## IN THE CLAIMS

- 1. (Currently amended) A method for detecting the presence of or predisposition to an ectodermal disorder comprising the steps of:
- (a) detecting the presence of a human TAJ gene or gene product in a cell of a host predetermined to be at elevated risk of having or being predisposed to a particular ectodermal disorder; and
- (b) correlating the presence of the TAJ gene or gene product with a presence of or predisposition to an the ectodermal disorder.
- 2. (Previously presented) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene.
- 3. (Previously presented) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript.
- 4. (Previously presented) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein.
- 5. (Previously presented) The method according to claim 1, wherein the detecting step is performed inferentially by determining a diagnostic sequence of the TAJ gene or gene product in the individual.
- 6. (Currently amended) The method according to claim 1, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to an the ectodermal disorder.
- 7. (Previously presented) The method according to claim 1, wherein the ectodermal disorder is an ectodermal dysplasia syndrome.
- 8. (Previously presented) The method according to claim 1, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 9-21. (Canceled)

- 22. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript, wherein the TAJ gene or gene product is truncated.
- 23. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein, wherein the TAJ gene or gene product is truncated.
- 24. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodennal disorder.
- 25. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder.
- 26. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder.
- 27. (New) The method according to claim 1, wherein the detecting step is performed inferentially by determining a diagnostic sequence of the TAJ gene or gene product in the individual, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder.
- 28. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript, wherein the TAJ gene or gene product is truncated, herein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder.
- 29. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein, wherein the TAJ gene or gene product is truncated, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder.

- 30. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 31. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, herein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 32. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 33. (New) The method according to claim 1, wherein the detecting step is performed inferentially by determining a diagnostic sequence of the TAJ gene or gene product in the individual, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 34. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ gene transcript, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.
- 35. (New) The method according to claim 1, wherein the detecting step comprises detecting a TAJ protein, wherein the TAJ gene or gene product is truncated, wherein the TAJ gene or gene product is a variant correlated with the presence of or predisposition to the ectodermal disorder, wherein the ectodermal disorder is an ectodermal dysplasia syndrome and the syndrome is Clouston syndrome.